

SEQUENCE LISTING

<110> Rondon, Isaac J
Ladner, Robert C

<120> BINDING PEPTIDES FOR CARCINOEMBRYONIC ANTIGEN (CEA)

<130> Sequence Listing DYX-016.1 US

<140> (not yet assigned)

<141> 2001-04-03

<150> US 09/541345

<151> 2000-04-03

<160> 151

<170> PatentIn Ver. 2.1

<210> 1

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding polypeptide

<220>

<221> VARIANT

<222> (1)

<223> Xaa is Asn, Asp or is absent

<220>

<221> VARIANT

<222> (2)

<223> Xaa is Trp

<220>

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<222> (3)

<223> Xaa is Asp, Phe or Val

<220>

<221> VARIANT

<222> (5)

<223> Xaa is Asn, Glu or Met

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 <222> (6)
 <223> Xaa is Asn, Leu, Met or Phe

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 <222> (16)

<223> Xaa is Leu, Ser, Trp or Tyr

<400> 1

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 2

<211> 16

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: family of
preferred CEA binding moieties

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<220>

<221> VARIANT

<222> (6)

<223> Xaa is Ph, Met, Leu or Asn

<220>

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<221> VARIANT

<222> (9)

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<222> (12)

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<220>

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<223> Xaa is Ala, Trp or Tyr

<221> VARIANT

<223> Xaa is Ala, Gly, His, Phe, Thr or Val

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys

5

10

<211> 16

<213> Artificial Sequence

<223> Description of Artificial Sequence: CEA binding polypeptide

Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser Tyr

1

5

10

15

<211> 16

<213> Artificial Sequence

<223> Description of Artificial Sequence: CEA binding polypeptide

Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu

1

5

10

15

<211> 16

<213> Artificial Sequence

<220>

Feedback

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<220>
<221> VARIANT
<222> (1)..(12)
<223> amino acid positions 4 and 9 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 2x10(8) different peptides
      based on the template sequence
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<210> 11
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<212> PRT
<213> Artificial Sequence
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<220>
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<222> (1)..(11)
<223> amino acid positions 3 and 9 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 1x10(9) different peptides
      based on the template sequence
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<210> 12
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[illegible]

<223> Description of Artificial Sequence: parental domain for design of microprotein display library

<221> VARIANT

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<223> amino acid positions 3 and 10 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 1x10(9) different peptides
      based on the template sequence
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Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
1 5 10

<211> 16

<213> Artificial Sequence

<223> Description of Artificial Sequence: parental domain for design of microprotein display library

<221> VARIANT

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<223> amino acid positions 4 and 13 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 2.5x10(8) different peptides
      based on the template sequence
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Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
1 5 10 15

<211> 16

<213> Artificial Sequence

8

<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

<220>

<221> VARIANT

<222> (1)..(3)

<223> Xaa is any amino acid except Cys

<220>

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<222> (5)..(6)

<223> Xaa is any amino acid except Cys

<400> 14

Xaa	Xaa	Xaa	Cys	Xaa	Xaa	Lys	Lys	Asp	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 15

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

<220>

<221> VARIANT

<222> (5)..(9)

<223> Xaa is any amino acid except Cys

<400> 15

Asp	Trp	Val	Cys	Xaa	Xaa	Xaa	Xaa	Xaa	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 16

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

<220>

<221> VARIANT

<222> (8)..(12)

<223> Xaa is any amino acid except Cys

<400> 16

Asp	Trp	Val	Cys	Glu	Asn	Lys	Xaa	Xaa	Xaa	Xaa	Xaa	Cys	Asn	Leu	Leu
1					5				10					15	

<210> 17

<211> 16

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: variable
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secondary library

<220>

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<222> (11)..(12)

<223> Xaa is any amino acid except Cys

<220>

<221> VARIANT

<222> (14)..(16)

<223> Xaa is any amino acid except Cys

<400> 17

Asp	Trp	Val	Cys	Glu	Asn	Lys	Lys	Asp	Gln	Xaa	Xaa	Cys	Xaa	Xaa	Xaa
1					5				10					15	

<210> 18

<211> 16

<212> PRT

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<223> Description of Artificial Sequence: variable
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secondary library

<220>

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<222> (6)..(7)

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<222> (9)

<223> Xaa is any amino acid except Cys

<220>

<221> VARIANT

<222> (12)

<223> Xaa is any amino acid except Cys

<220>

<221> VARIANT

<222> (15)

<223> Xaa is any amino acid except Cys

<400> 18

Asp	Trp	Val	Cys	Glu	Xaa	Xaa	Lys	Xaa	Gln	Trp	Xaa	Cys	Asn	Xaa	Leu
1				5					10					15	

<210> 19

<211> 16

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<222> (5)..(7)

<223> Xaa is any amino acid except Cys

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<222> (9)

<223> Xaa is any amino acid except Cys

<220>

<221> VARIANT

<222> (12)

<223> Xaa is any amino acid except Cys

<400> 19

Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
1 5 10 15

<210> 20

<211> 16

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<223> Description of Artificial Sequence: variable
sublibrary sequence used in designing focused
secondary library

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<222> (1)

<223> Xaa is any amino acid except Cys

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<222> (3)

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<221> VARIANT

<222> (14)..(16)

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<400> 20

Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
1 5 10 15

<210> 21

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: isolate of
TN10/9 library found not to bind CEA

<400> 21

Asn Trp Arg Cys Lys Leu Phe Pro Arg Tyr Pro Tyr Cys Ser Ser Trp
1 5 10 15

DEFEAT

```
<400> 22
Arg Tyr Cys Glu Phe Phe Pro Trp Ser Leu His Cys Gly Arg Pro
  1             5             10             15
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<220>
<223> Description of Artificial Sequence: conserved
      amino acid positions in first family of CEA
      binding peptides
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<220>
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<223> X is Asn, Leu, Met or Phe
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<220>
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<223> X is Asp, Gly, Ile, Lys, Phe or Thr

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<220>
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<222> (9)
<223> X is Arg, Asn, Asp, Glu or Gly
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<220>
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<222> (12)
<223> X is Ala, Gly, His, Phe, Thr or Val
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13

<223> X is Arg, Leu, Pro or Ser

<400> 23

Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
1 5 10 15

<210> 24

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 24

Ser Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser
1 5 10 15

Tyr Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
20 25

<210> 25

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 25

Ser Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu
1 5 10 15

Leu Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
20 25

<210> 26

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 26

Ser Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro
1 5 10 15

Trp Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
20 25

<210> 27

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic CEA
binding peptide with C-terminal immobilization
sequence

<400> 27

Ser Asp Trp Val Cys Glu Leu Thr Thr Gly Gly Tyr Val Cys Gln Pro
1 5 10 15

Leu Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
20 25

<210> 28

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: C-terminal
sequence for immobilizing peptides

<400> 28

Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
1 5 10

<210> 29

<211> 16

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

<220>

<221> VARIANT

<222> (1)..(3)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (5)..(6)

<223> X is any amino acid except Cys

<400> 29

Xaa	Xaa	Xaa	Cys	Xaa	Xaa	Lys	Lys	Asp	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 30

<211> 16

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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

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<222> (5)..(9)

<223> X is any amino acid except Cys

<400> 30

Asp	Trp	Val	Cys	Xaa	Xaa	Xaa	Xaa	Xaa	Gln	Trp	Thr	Cys	Asn	Leu	Leu
1				5					10					15	

<210> 31

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

<220>

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<222> (8)..(12)

<223> X is any amino acid except Cys

<400> 31

Asp Trp Val Cys Glu Asn Lys Xaa Xaa Xaa Xaa Xaa Cys Asn Leu Leu
1 5 10 15

<210> 32

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

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<222> (11)..(12)

<223> X is any amino acid except Cys

<220>

<221> VARIANT

<222> (14)..(16)

<223> X is any amino acid except Cys

<400> 32

Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Xaa Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 33

<211> 16

<212> PRT

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<223> Description of Artificial Sequence: template
sequence for sublibrary used in construction of
focused secondary display library

FTS2200

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<223> X is any amino acid except Cys

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<223> X is any amino acid except Cys

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<400> 33
Asp Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
1 5 10 15

<210> 34
<211> 16
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sequence for sublibrary used in construction of
focused secondary display library

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<223> X is any amino acid except Cys

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<222> (12)

<223> X is any amino acid except Cys

<400> 34

Asn Trp Val Cys Xaa Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Ser Tyr
1 5 10 15

<210> 35

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: template
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focused secondary display library

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<222> (1)

<223> X is any amino acid except Cys

<220>

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<222> (3)

<223> X is any amino acid except Cys

<220>

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<222> (14)..(16)

<223> X is any amino acid except Cys

<400> 35

Xaa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
1 5 10 15

<210> 36

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: family of CEA
binding polypeptides

<220>

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<222> (1)

<223> Xaa is Asp, Asn, Ala or Ile

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<221> VARIANT

<222> (3)

<223> Xaa is Val, Ile, Met, Tyr, Phe, Pro or Asp

<220>

<221> VARIANT

<222> (5)

<223> Xaa is Asn, Glu or Asp

<220>

<221> VARIANT

<222> (6)

<223> Xaa is Leu, Phe, Tyr, Trp, Val Met, Ile or Asn

<220>

<221> VARIANT

<222> (7)

<223> Xaa is Phe, Leu, Asp, Glu, Ala, Ile, Lys, Asn,
Ser, Val, Trp or Tyr

<220>

<221> VARIANT

<222> (8)

<223> Xaa is Lys, Phe, Asp, Gly, Leu, Asn or Trp

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<221> VARIANT

<222> (9)

<223> Xaa is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gln
or Trp

<220>

<221> VARIANT

<222> (10)

<223> Xaa is Gln or Lys

<220>

<221> VARIANT

<222> (12)

<223> Xaa is Phe, Thr, Met, Ser, Ala, Asn, Val, His,
Ile, Pro, Trp or Tyr

<220>

<221> VARIANT

<222> (14)

<223> Xaa is Asn, Asp, Glu, Pro, Gln or Ser

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<221> VARIANT

<222> (15)

<223> Xaa is Val, Leu, Ile, Pro, Ala, Gln, Ser, Met,
Glu,Thr, Lys or Trp

<220>

<221> VARIANT

<222> (16)

<223> Xaa is Leu, Met, Val, Tyr, Ala, Ile, Trp, His,
Pro, Gln, Glu, Phe, Lys or Arg

<400> 36

Xaa Trp Xaa Cys Xaa Xaa Xaa Xaa Xaa Trp Xaa Cys Xaa Xaa Xaa
1 5 10 15

<210> 37

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 37

Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
1 5 10 15

<210> 38

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 38

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Met
1 5 10 15

<210> 39
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 39
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Met
1 5 10 15

<210> 40
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 40
Asn Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Gln Glu
1 5 10 15

<210> 41
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CEA binding
polypeptide

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Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Gln Val Lys
1 5 10 15

<210> 42
<211> 16
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1

5

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15

<210> 46

<211> 16

<212> PRT

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polypeptide

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Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Val
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<210> 47

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

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Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ala Ile
1 5 10 15

<210> 48

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 48

Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Met Ala
1 5 10 15

<210> 49

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 49

Asp	Trp	Val	Cys	Glu	Phe	Leu	Lys	Met	Gln	Trp	Ala	Cys	Asn	Val	Leu
1					5				10					15	

<210> 50

<211> 16

<212> PRT

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<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 50

Asp	Trp	Val	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Asn	Val	Met
1				5					10					15	

<210> 51

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 51

Ala	Trp	Pro	Cys	Asn	Leu	Phe	Lys	Asn	Gln	Trp	Phe	Cys	Pro	Pro	Gln
1				5					10					15	

<210> 52

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 52

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Leu
1 5 10 15

<210> 53

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 53

Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Lys Trp
1 5 10 15

<210> 54

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 54

Asp Trp Val Cys Glu Trp Leu Lys Met Gln Trp Ala Cys Asn Met Leu
1 5 10 15

<210> 55

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CEA binding
polypeptide

<400> 55

Asp Trp Val Cys Asp Phe Phe Phe Asn Gln Trp Thr Cys Asn Leu Leu
1 5 10 15

<210> 56

Variable	Mean	SD	Min	Max
Age	34.5	10.2	21	55
Gender	Male	Female		
Marital status	Married	Single		
Education	High school	College		
Occupation	Manager	Worker		
Income	Low	High		
Health status	Good	Poor		
Stress level	Low	High		
Life satisfaction	Low	High		
Depression	Low	High		
Loneliness	Low	High		
Self-esteem	Low	High		
Resilience	Low	High		
Optimism	Low	High		
Gratitude	Low	High		
Forgiveness	Low	High		
Empathy	Low	High		
Compassion	Low	High		
Kindness	Low	High		
Generosity	Low	High		
Patience	Low	High		
Humility	Low	High		
Modesty	Low	High		
Shyness	Low	High		
Introversion	Low	High		
Extroversion	Low	High		
Sensitivity	Low	High		
Emotionality	Low	High		
Impulsivity	Low	High		
Risk-taking	Low	High		
Curiosity	Low	High		
Openness	Low	High		
Conscientiousness	Low	High		
Agreeableness	Low	High		
Neuroticism	Low	High		
Stability	Low	High		
Control	Low	High		
Power	Low	High		
Authority	Low	High		
Leadership	Low	High		
Influence	Low	High		
Impact	Low	High		
Significance	Low	High		
Importance	Low	High		
Value	Low	High		
Meaning	Low	High		
Purpose	Low	High		
Direction	Low	High		
Focus	Low	High		
Attention	Low	High		
Interest	Low	High		
Engagement	Low	High		
Participation	Low	High		
Involvement	Low	High		
Commitment	Low	High		
Dedication	Low	High		
Devotion	Low	High		
Loyalty	Low	High		
Fidelity	Low	High		
Trustworthiness	Low	High		
Reliability	Low	High		
Consistency	Low	High		
Stability	Low	High		
Endurance	Low	High		
Persistence	Low	High		
Perseverance	Low	High		
Fortitude	Low	High		
Steadfastness	Low	High		
Unwaveringness	Low	High		
Immutability	Low	High		
Indestructibility	Low	High		
Invulnerability	Low	High		
Impenetrability	Low	High		
Impassability	Low	High		
Impermeability	Low	High		
Immutability	Low	High		
Indestructibility	Low	High		
Invulnerability	Low	High		
Impenetrability	Low	High		
Impassability	Low	High		
Impermeability	Low	High		

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Dedication

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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<210> 126

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16-mer microprotein analogues

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<210> 127

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16-mer microprotein analogues

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<210> 128

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<400> 128

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

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16-mer microprotein analogues

<400> 137

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